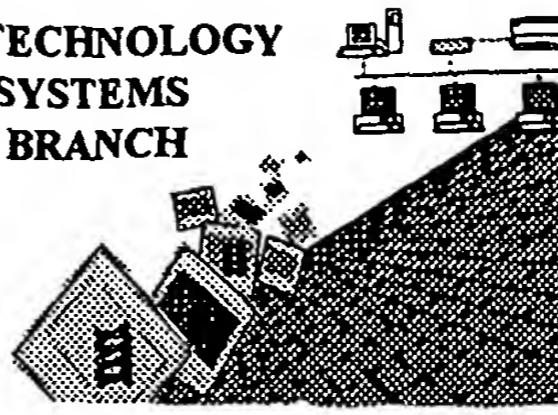


87-100

BIOTECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/053,645
Source: OIPE
Date Processed by STIC: 2/7/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name,
Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two,
2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office,
Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

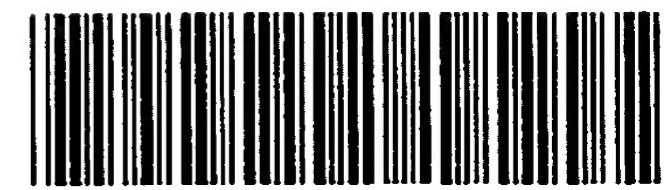
Raw Sequence Listing Error Summary

<u>ERROR DETECTED</u>	<u>SUGGESTED CORRECTION</u>	<u>SERIAL NUMBER:</u> <u>10/053,645</u>
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ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleic
Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
"bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
(OLD RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
(NEW RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 9 Use of n's or Xaa's
(NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <213>
Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
1-19 (and more)
- 11 Use of <220> Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
"bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.



OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/053,645

DATE: 02/07/2002
TIME: 12:30:12

Input Set : A:\SEQ LISTING OF 10412-022.TXT
Output Set: N:\CRF3\02072002\J053645.raw

Does Not Comply
Corrected Diskette Needed

4 <110> APPLICANT: Robert E. Klem
 6 <120> TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TREATING A
 7 CELL-PROLIFERATIVE DISORDER USING CRE DECOY OLIGOMERS, BCL-2
 8 ANTISENSE OLIGOMERS, AND HYBRID OLIGOMERS THEREOF
 11 <130> FILE REFERENCE: 10412-022-999
 12 <140> CURRENT APPLICATION NUMBER: US/10/053,645 *PR 14*
 13 <141> CURRENT FILING DATE: 2002-01-22
 16 <150> PRIOR APPLICATION NUMBER: 60/263,244
 17 <151> PRIOR FILING DATE: 2001-01-22
 19 <160> NUMBER OF SEQ ID NOS: 43
 21 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 23 <210> SEQ ID NO: 1
 24 <211> LENGTH: 20 *Artificial (global misspellng)*
 25 <212> TYPE: DNA
 C--> 26 <213> ORGANISM: *Artificail* sequence → see item 1 on Error Summary
 W--> 28 <220> FEATURE:
 W--> 28 <223> OTHER INFORMATION:
 28 <400> SEQUENCE: 1
 29 cagcgtgcgc catccttccc
 31 <210> SEQ ID NO: 2
 32 <211> LENGTH: 35
 33 <212> TYPE: DNA
 C--> 34 <213> ORGANISM: *Artificail* sequence 20
 W--> 36 <220> FEATURE:
 W--> 36 <223> OTHER INFORMATION:
 36 <400> SEQUENCE: 2
 37 ctttcctct ggaaaggatg ggcacgctg ggaga 35
 39 <210> SEQ ID NO: 3
 40 <211> LENGTH: 20
 41 <212> TYPE: DNA
 C--> 42 <213> ORGANISM: *Artificail* sequence
 W--> 44 <220> FEATURE:
 W--> 44 <223> OTHER INFORMATION:
 44 <400> SEQUENCE: 3
 45 gatgcaccta cccagcctcc 20
 47 <210> SEQ ID NO: 4
 48 <211> LENGTH: 33
 49 <212> TYPE: DNA
 C--> 50 <213> ORGANISM: *Artificail* sequence
 W--> 52 <220> FEATURE:
 W--> 52 <223> OTHER INFORMATION:
 52 <400> SEQUENCE: 4
 53 acgggtacg gaggtgggt aggtgcattt ggt 33

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/053,645

DATE: 02/07/2002
TIME: 12:30:12

Input Set : A:\SEQ LISTING OF 10412-022.TXT
Output Set: N:\CRF3\02072002\J053645.raw

```

55 <210> SEQ ID NO: 5
56 <211> LENGTH: 20
57 <212> TYPE: DNA
C--> 58 <213> ORGANISM: Artificail sequence
W--> 60 <220> FEATURE:
W--> 60 <223> OTHER INFORMATION:
60 <400> SEQUENCE: 5
61 acaaaggcat cctgcagttg 20
63 <210> SEQ ID NO: 6
64 <211> LENGTH: 36
65 <212> TYPE: DNA
C--> 66 <213> ORGANISM: Artificail sequence
W--> 68 <220> FEATURE:
W--> 68 <223> OTHER INFORMATION:
68 <400> SEQUENCE: 6
69 cccccaactg caggatgcct ttgtggaact gtacgg 36
71 <210> SEQ ID NO: 7
72 <211> LENGTH: 20
73 <212> TYPE: DNA
C--> 74 <213> ORGANISM: Artificail sequence
W--> 76 <220> FEATURE:
W--> 76 <223> OTHER INFORMATION:
76 <400> SEQUENCE: 7
77 gggaaaggatg ggcacgctg 20
79 <210> SEQ ID NO: 8
80 <211> LENGTH: 17
81 <212> TYPE: DNA
C--> 82 <213> ORGANISM: Artificail sequence
W--> 84 <220> FEATURE:
W--> 84 <223> OTHER INFORMATION:
84 <400> SEQUENCE: 8
85 cgctgtgcac cctcttg 17
87 <210> SEQ ID NO: 9
88 <211> LENGTH: 17
89 <212> TYPE: DNA
C--> 90 <213> ORGANISM: Artificail sequence
W--> 92 <220> FEATURE:
W--> 92 <223> OTHER INFORMATION:
92 <400> SEQUENCE: 9
93 taccgcgtgc gaccctc 17
95 <210> SEQ ID NO: 10
96 <211> LENGTH: 17
97 <212> TYPE: DNA
C--> 98 <213> ORGANISM: Artificail sequence
W--> 100 <220> FEATURE:
W--> 100 <223> OTHER INFORMATION:
100 <400> SEQUENCE: 10
101 tccttaccgcg tgcgacc 17
103 <210> SEQ ID NO: 11

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/053,645

DATE: 02/07/2002
TIME: 12:30:12

Input Set : A:\SEQ LISTING OF 10412-022.TXT
Output Set: N:\CRF3\02072002\J053645.raw

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104 <211> LENGTH: 17
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C--> 106 <213> ORGANISM: Artificail sequence
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  109 ctttcctacc gcgtgcg
  111 <210> SEQ ID NO: 12
  112 <211> LENGTH: 17
  113 <212> TYPE: DNA
C--> 114 <213> ORGANISM: Artificail sequence
W--> 116 <220> FEATURE:
W--> 116 <223> OTHER INFORMATION:
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  117 gacccttcct accgcgt
  119 <210> SEQ ID NO: 13
  120 <211> LENGTH: 17
  121 <212> TYPE: DNA
C--> 122 <213> ORGANISM: Artificail sequence
W--> 124 <220> FEATURE:
W--> 124 <223> OTHER INFORMATION:
  124 <400> SEQUENCE: 13
  125 ggagaccctt cctaccg
  127 <210> SEQ ID NO: 14
  128 <211> LENGTH: 15
  129 <212> TYPE: DNA
C--> 130 <213> ORGANISM: Artificail sequence
W--> 132 <220> FEATURE:
W--> 132 <223> OTHER INFORMATION:
  132 <400> SEQUENCE: 14
  133 gggcgccgac cgccgg
  135 <210> SEQ ID NO: 15
  136 <211> LENGTH: 15
  137 <212> TYPE: DNA
C--> 138 <213> ORGANISM: Artificail sequence
W--> 140 <220> FEATURE:
W--> 140 <223> OTHER INFORMATION:
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  141 cggcggggcg acgga
  143 <210> SEQ ID NO: 16
  144 <211> LENGTH: 16
  145 <212> TYPE: DNA
C--> 146 <213> ORGANISM: Artificail sequence
W--> 148 <220> FEATURE:
W--> 148 <223> OTHER INFORMATION:
  148 <400> SEQUENCE: 16
  149 cgggagcgcg gcgggc
  151 <210> SEQ ID NO: 17
  152 <211> LENGTH: 18

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/053,645

DATE: 02/07/2002
TIME: 12:30:12

Input Set : A:\SEQ LISTING OF 10412-022.TXT
Output Set: N:\CRF3\02072002\J053645.raw

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153 <212> TYPE: DNA
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W--> 156 <220> FEATURE:
W--> 156 <223> OTHER INFORMATION:
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157 tctcccagcg tgcgccat
159 <210> SEQ ID NO: 18
160 <211> LENGTH: 18
161 <212> TYPE: DNA
C--> 162 <213> ORGANISM: Artificail sequence
W--> 164 <220> FEATURE:
W--> 164 <223> OTHER INFORMATION:
164 <400> SEQUENCE: 18
165 tgcactcacg ctcggcct
167 <210> SEQ ID NO: 19
168 <211> LENGTH: 106
169 <212> TYPE: DNA
C--> 170 <213> ORGANISM: Artificail sequence
W--> 172 <220> FEATURE:
W--> 172 <223> OTHER INFORMATION:
172 <400> SEQUENCE: 19
173 gcgcgcgcgc ctccgcgcgc cctgcgcgcgc gccccgcgc gctccgcgc gccgcgtctcc 60
174 ccttattgtt aaaaacatgt tagaagcaat gaatgtatat aaaagc 106
176 <210> SEQ ID NO: 20
177 <211> LENGTH: 717
178 <212> TYPE: DNA
179 <213> ORGANISM: Homo Sapiens
181 <220> FEATURE:
182 <221> NAME/KEY: CDS
183 <222> LOCATION: (1)...(717)
185 <400> SEQUENCE: 20
186 atg gcg cac gct ggg aga acg ggg tac gac aac cgg gag ata gtg atg 48
187 Met Ala His Ala Gly Arg Thr Gly Tyr Asp Asn Arg Glu Ile Val Met
188 1 5 10 15
190 aag tac atc cat tat aag ctg tcg cag agg ggc tac gag tgg gat gcg 96
191 Lys Tyr Ile His Tyr Lys Leu Ser Gln Arg Gly Tyr Glu Trp Asp Ala
192 20 25 30
194 gga gat gtg ggc gcc gcg ccc ccg ggg gcc ccc gca ccg ggc atc 144
195 Gly Asp Val Gly Ala Ala Pro Pro Gly Ala Ala Pro Ala Pro Gly Ile
196 35 40 45
198 ttc tcc tcc cag ccc ggg cac acg ccc cat cca gcc gca tcc cgc gac 192
199 Phe Ser Ser Gln Pro Gly His Thr Pro His Pro Ala Ala Ser Arg Asp
200 50 55 60
202 ccg gtc gcc agg acc tcg ccg ctg cag acc ccg gct gcc ccc ggc gcc 240
203 Pro Val Ala Arg Thr Ser Pro Leu Gln Thr Pro Ala Ala Pro Gly Ala
204 65 70 75 80
206 gcc gcg ggg cct gcg ctc agc ccg gtg cca cct gtg gtc cac ctg gcc 288
207 Ala Ala Gly Pro Ala Leu Ser Pro Val Pro Pro Val Val His Leu Ala
208 85 90 95

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The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/053,645

DATE: 02/07/2002
TIME: 12:30:12

Input Set : A:\SEQ LISTING OF 10412-022.TXT
Output Set: N:\CRF3\02072002\J053645.raw

210	ctc	cgc	caa	gcc	ggc	gac	gac	ttc	tcc	cgc	cgc	tac	cgc	ggc	gac	ttc	336	
211	Leu	Arg	Gln	Ala	Gly	Asp	Asp	Phe	Ser	Arg	Arg	Tyr	Arg	Gly	Asp	Phe		
212				100				105								110		
214	gcc	gag	atg	tcc	agc	cag	ctg	cac	ctg	acg	ccc	ttc	acc	gcg	cg	gga	384	
215	Ala	Glu	Met	Ser	Ser	Gln	Leu	His	Leu	Thr	Pro	Phe	Thr	Ala	Arg	Gly		
216				115				120								125		
218	cgc	ttt	gcc	acg	gtg	gtg	gag	gag	ctc	ttc	agg	gac	ggg	gtg	aac	tgg	432	
219	Arg	Phe	Ala	Thr	Val	Val	Glu	Glu	Leu	Phe	Arg	Asp	Gly	Val	Asn	Trp		
220				130				135								140		
222	ggg	agg	att	gtg	gcc	ttc	ttt	gag	ttc	ggt	ggg	gtc	atg	tgt	gtg	gag	480	
223	Gly	Arg	Ile	Val	Ala	Phe	Phe	Glu	Phe	Gly	Gly	Val	Met	Cys	Val	Glu		
224				145				150								160		
226	agc	gtc	aac	cgg	gag	atg	tcg	ccc	ctg	gtg	gac	aac	atc	gcc	ctg	tgg	528	
227	Ser	Val	Asn	Arg	Glu	Met	Ser	Pro	Leu	Val	Asp	Asn	Ile	Ala	Leu	Trp		
228				165				170								175		
230	atg	act	gag	tac	ctg	aac	cgg	cac	ctg	cac	acc	tgg	atc	cag	gat	aac	576	
231	Met	Thr	Glu	Tyr	Leu	Asn	Arg	His	Leu	His	Thr	Trp	Ile	Gln	Asp	Asn		
232				180				185								190		
234	gga	ggc	tgg	gat	gcc	ttt	gtg	gaa	ctg	tac	ggc	ccc	agc	atg	cg	cct	624	
235	Gly	Gly	Trp	Asp	Ala	Phe	Val	Glu	Leu	Tyr	Gly	Pro	Ser	Met	Arg	Pro		
236				195				200								205		
238	ctg	ttt	gat	ttc	tcc	tgg	ctg	tct	ctg	aag	act	ctg	ctc	agt	ttg	gcc	672	
239	Leu	Phe	Asp	Phe	Ser	Trp	Leu	Ser	Leu	Lys	Thr	Leu	Leu	Ser	Leu	Ala		
240				210				215								220		
242	ctg	gtg	gga	gct	tgc	atc	acc	ctg	ggt	gcc	tat	ctg	agc	cac	aag		717	
243	Leu	Val	Gly	Ala	Cys	Ile	Thr	Leu	Gly	Ala	Tyr	Leu	Ser	His	Lys			
244				225				230								235		
248	<210>	SEQ	ID	NO:	21													
249	<211>	LENGTH:	239															
250	<212>	TYPE:	PRT															
251	<213>	ORGANISM:	Homo Sapiens															
253	<400>	SEQUENCE:	21															
254	Met	Ala	His	Ala	Gly	Arg	Thr	Gly	Tyr	Asp	Asn	Arg	Glu	Ile	Val	Met		
255				1				5								15		
256	Lys	Tyr	Ile	His	Tyr	Lys	Leu	Ser	Gln	Arg	Gly	Tyr	Glu	Trp	Asp	Ala		
257					20				25							30		
258	Gly	Asp	Val	Gly	Ala	Ala	Pro	Pro	Gly	Ala	Ala	Pro	Ala	Pro	Gly	Ile		
259					35				40							45		
260	Phe	Ser	Ser	Gln	Pro	Gly	His	Thr	Pro	His	Pro	Ala	Ala	Ser	Arg	Asp		
261					50				55							60		
262	Pro	Val	Ala	Arg	Thr	Ser	Pro	Leu	Gln	Thr	Pro	Ala	Ala	Pro	Gly	Ala		
263				65				70								80		
264	Ala	Ala	Gly	Pro	Ala	Leu	Ser	Pro	Val	Pro	Pro	Val	Val	His	Leu	Ala		
265					85				90							95		
266	Leu	Arg	Gln	Ala	Gly	Asp	Asp	Phe	Ser	Arg	Arg	Tyr	Arg	Gly	Asp	Phe		
267					100				105							110		
268	Ala	Glu	Met	Ser	Ser	Gln	Leu	His	Leu	Thr	Pro	Phe	Thr	Ala	Arg	Gly		
269					115				120							125		
270	Arg	Phe	Ala	Thr	Val	Val	Glu	Glu	Leu	Phe	Arg	Asp	Gly	Val	Asn	Trp		

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/053,645

DATE: 02/07/2002
TIME: 12:30:13

Input Set : A:\SEQ LISTING OF 10412-022.TXT
Output Set: N:\CRF3\02072002\J053645.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application Number
L:26 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
L:28 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:28 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:34 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2
L:36 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:36 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:42 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:44 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:44 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:50 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:52 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:52 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:58 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:60 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:60 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:66 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
L:68 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:68 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:74 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
L:76 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:76 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:82 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
L:84 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:84 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:90 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
L:92 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:92 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:98 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
L:100 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:100 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:106 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
L:108 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:108 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:114 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12
L:116 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:116 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:122 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:13
L:124 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:124 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:130 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
L:132 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:132 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:138 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15
L:140 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:140 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:146 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16
L:148 M:258 W: Mandatory Feature missing, <220> FEATURE:

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/053,645

DATE: 02/07/2002
TIME: 12:30:13

Input Set : A:\SEQ LISTING OF 10412-022.TXT
Output Set: N:\CRF3\02072002\J053645.raw

L:148 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:154 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:17
L:156 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:156 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:162 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:18
L:164 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:164 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:170 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:19
L:172 M:258 W: Mandatory Feature missing, <220> FEATURE:
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L:388 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
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L:412 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
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L:428 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:428 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: